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## IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of

Ivarie et all.

Application No: 09/173,864

Filed:

October 16, 1998

: Examiner:

: Group Art Unit: 1633

Sumesh Kaushai

Title:

NOVEL TRANSGENIC BIRDS AND THEIR EGGS

Assistant Commissioner for Patents

Washington, DC 20231

Sir:

## DECLARATION UNDER 37 CFR 1.132

- I, Jeffrey Rapp, Ph.D. hereby declare as follows:
- 1. I am a staff scientist employed by AviGenics, inc., of Athens, GA., licensec of the subject application, since June 1999. I consider myself to be skilled in the art of avian transgenesis. My curriculum vitae is appended hereto.

- 2. The following reports experiments conducted by me personally, or by those working under my supervision, to produce a transgenic chieken expressing an exogenous protein, following the procedures set forth in the subject application of Ivarie, e al.
- One set of experiments was designed to demonstrate the expression of human interferon in a chicken, as suggested at page 32, line 6 of the specification.
- 4. Following the teachings of Example 1 (Vector Construction) of the specification, an pNLB-CMV-IFN vector was created, substituting an IFN encoding sequence for the BL encoding sequence of the Example.

The DNA sequence for human interfetion α-2b based on hen oviduet optimized codon usage was created using the BACKTRANSLATE program of the Wisconsin Package, version 9.1 (Genetics Computer Group. Inc., Madison, WI) with a codon usage table compiled from the chicken (Gallus gallus) ovalbumin, lysozyme, ovomucoid, and ovotransferrin proteins. The template and primer oligonucleotides listed in Table 1 were amplified by PCR with Pfu polymerase (Stratagene, La Jolla, CA) using 20 cycles of 94 °C for 1 min., 50°C for 30 sec., and 72°C for 1 min. and 10 sec. PCR products were purified from a 12% polyacrylamide-TBE gel by the "crush and soak" method (Maniatis et al. 1982), then combined as templates in an amplification reaction using only IFN-1 and IFN-8 as primers. The resulting PCR product was digested with Hind III and Xba I and gel purified from a 2% agarose-TAE gel, then ligated into Hind III and Xba I digested, alkaline phosphatase-treated pBluescript KS (Statagene), resulting in the plastmid pBluKSP-IFNMagMax. Both strands were sequenced by cycle sequencing on an ABI PRISM 377 DNA Sequencer (Perkin-Elmer, Foster City, CA) using

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Declaration Jeffrey Rapp, Ph.D.

universal T7 or T3 primers. Mutations in pBluKSP-IFN derived from the original oligonucleotide templates were corrected by site-directed mulagenesis with the Transformer Site-Directed Mutagenesis Kit (Clontech, Palo Alto, CA). The IFN coding sequence was then removed from the corrected pBluKSP-IFN with Hind III and Xba 1, purified from a 0.8% agarose-TAE Gel, and ligated to Hind III and Xba I digested, alkaline phosphatase-treated pCMV-BetaLa-3B-dH. The resulting plasmid was pCMV-IFN which contained IFN coding sequence controlled by the cytomegalovirus immediate early promoter/enhancer and SV40 polyA site. To clone the IFN coding sequence controlled by the CMV promoter/enhancer into the NLB retroviral plasmid, pCMV-IFN was first digested with ClaI and XbaI, then both ends were filled in with klenow fragment of DNA polymerase (New England BioLabs, Beverly, MA), pNLB-adapter was digested with Ndel and Kpnl, and both ends were made blunt by T4 DNA polymerase (New England BioLabs). Appropriate DNA fragments were purified on a 0.8% agarose-TAE get, then ligated and transformed into D#5α cells. The resulting plasmid was pNLB-adapter-CMV-IFN. This plasmid was then digested with MluI and partially digested with BlpI and the appropriate fragment was gel purified. pNLB-CMV-EGFP was digested with Mlul and Blpl, then alkaline-phosphatase recated and gel purified. The Mlul/Blpl partial fragment of pNLB-adapter-CMV-IFN was ligated to the large fragment derived from the Mlul/BlpI digest of nNLB-CMV-EGFP, creating pNLB-CMV-IFN.

5. Following the procedures of Example 2 (Production of Transduction Particles), transduction particles of pNLB-CMV-IFN were produced

Senta packaging cells (Cosset et al., 1990) were plated at a density of 3 x 105 cells/35mm tissue culture dish in F-10 medium (Life Technologies) supplemented with 50%

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calf serum (Atlanta Biologicals), 1% chicken serum (Life Technologies), 50 μg/ml hygromycin (Sigma), and 50 µg/ml phleomycin (CAYLA, Toulouse, France). These cells were transfected 24h after plating with 2 μg of CsCl-purified pNLB-CMV-IFN DNA and 6 μl of Lipofectin liposomes (Life Technologies) in a final volume of 500 µl Optimem (Life Technologies). The plates were gently rocked for four hours at 37 C in a 5% CO incubator. For each well, the media was removed, washed once with 1 ml of Optimern and re-fed with 2 mls of F-10 medium supplemented with 50% calf scrum, 1% chicken serum, 50 µg/ml hygromycin, and 50 µg/ml phleomycin. The next day, medium from transfected Sentas was recovered and filtered through a 0.45 micron filter. This medium was then used to transduce Isolde cells. 0.3 ml of the filtered medium recovered from Senta cells was added to 9.6 ml of F-10 (Life Technologies) supplemented as described above, in addition to polybrene (Sigma) at a final concentration of 4 µg/ml. This mixture was added to 106 Isolde packaging cells (Cosset et al., 1990) plated on a 100ram dish the previous day, then replaced with fresh F-10 medium (as described for Senta growth) 4h later. The next day, the medium was replaced with fresh medium which also contained 200 µg/ml neomycin (G418, Sigma). Every other day, the medium was replaced with fresh F- 10 medium supplemented with 50% calf serum, 1% chicken serum, 50 kg/ml hygromycin, 50 kg/ml phleomycin, and 200 µg/ml neomycin. Eleven to twelve days later, single colonies were visible by eye, and these were picked and placed into 24 well dishes. When some of the 24 well dishes became confluent, medium was harvested and titered to determine the cell lines with the highest production of retrovirus. Titering was performed by plating 7.5 x 10<sup>4</sup> Senta cells per well in 24 well plates on the day prior to viral harvest and transduction. The next day 1.0 ml of fresh F-10 medium

supplemented with 50% calf serum, 1% chicken serum, 50 µg/ml hygromycin, and 50 µg/ml phleomycin was added to each well of the isolated Isolde colonies. Virus was harvested for 8-10 hours. The relative density of each well of Isoldes was noted. After 8-10 hours, 2 and 20µl of media from each well of Isoldes was added directly to the media of duplicate wills of the Sentas. Harvested medium was also tested for the presence of IFN by IFN ELISA and for IFN bioreactivity. The next day the media was replaced with F- 10 medium supplemented with 50% calf serum. 1% chicken serum, 50 µg/ml hygromycin, 50 µg/ml phleomycin, and 200 µg/ml neomycin. When obvious neomycin resistant colonies were evident in the wells of transduced Sentas, the number of colonies was counted for each well. The Isolde colony producing the highest titer was determined by taking into account the number of colonies and correcting for the density of the Isolde cells when the viral particles were harvested (i.e., if two Isolde colonies gave rise to media with the same titer, but one was at a 5% density abd the other was at a 50% density at the time of viral harvest, the one at the 5% density was chosen for further work, as was the case in the present example).

The Isolde cell fine producing the highest titer of IFN-encoding transducing particles was scaled up to six T 75 tissue culture flasks. When flasks were confluent, cells were washed with F-10 medium (unsupplemented) and transducing particles were then harvested for I6h in 14 ml/flask of F-10 containing 1% calf serum (Atlanta Biologicals) and 0.2% chicken serum (Life Technolocyies). Medium was harvested, filtered through a 0.45 micron syringe filter, then centrifuged at 195,000xg in a Beckman 60Ti rotor for 35 min. Liquid was removed except for 1 ml, and this was incubated with the pellet at 37°C with gentle shaking for one hour. Aliquots were frozen at -70°C. Transducing particles were then titered on Senta cells to determine concentrations used to inject embryos.

Following the procedures of Example 3 Production of Transgenic Chickens), chimeric birds were produced.

Approximately 300 White Leghorn (strain Line 0) eggs were windowed according to the Speksnijder procedure (U.S. Patent No. 5,897,998), then injected with ~7 x 104 transducing particles per egg. Eggs hatched 21 days after injection, and human IFN levels were measured by IFN ELISA from serum samples collected from chicks one week after hatch.

Following the procedures of Example 10 (Production of Fully Transgenic G1 Chickens), males were selected for breeding.

To screen for to roosters which contained the IFN transgene in their sperm, DNA was extracted from rooster sperm samples by Chelex-100 extraction (Walsh et al., 1991). DNA samples were then subjected to Taqman analysis on a 7700 Sequence Detector (Perkin Elmer) using the "neo tor-1" (5'-TGGATTGCA@GCAGGTTCT-3') and "neo rev-1" (5'-GTGCCCAGTCATAGCCGAAT-3') primers and FAM labeled NEO-PROBE1 (5'-CCTCTCCACCCAAGCGGCCG-3') to detect the transgene. Three Go roosters with the highest levels of the transgene in their sperm samples were bred to nontransgenic SPAFAS (White Leghorn) hens by artificial insemination.

Blood DNA samples were screened for the presence of the transgene by Taqman™ analysis as described above. Out of 1,597 offspring, one rooter was found to be transgenic (a.k.a. "Alphie"). Alphie's serum was tested for the presence of hIFN by hIFN ELISA, and hIFN was present at 200 nanograms/ml.

Alphie's sperre was used for artificial the mination of nontransgenic SPAFAS (White Leghorn) hens. To date 106 out of 202 (~52%) offspring contain the transgene as detected by Tagman™ analysis. These breeding results follow a Mendelian inheritance pattern and indicate that Alphie is transgeric

Alphie was maked with nontransgenic White Leghorn hens to produce G₂ offspring, and blood DNA from G2 chicks was assayed by Tagman analysis as described above. Of the 245 chicks tested, 149 (6018%) contained the transgene, indicating Mendelian inheritance of the transgene. Egg white material from 150 of the G2 hens has been assayed to date for hIFN by ELISA and found to contain an average of about 900 nanograms/ml hIFN. G2 hens that did not contain the transgene, as determined by Tadman analysis, also did not contain hIFN in their egg white.

- In other experiments, additional proteins were produced in transgenic chickens, including 8. GM-CSF, human erythropoietin (EPO), and \(\beta\)-lactamase (\(\beta\)-lac, as suggested on page 32, lines 3-13, of the subject application of Ivarie, et al. These additional proteins were produced using the procedures as set forth above (pNLB-CMV-GMCSF and pNLB-CMV- β-lac vectors), except that the promoter used for the production of EPO was a synthetic promoter, MDOT, constructed as suggested on page 11, lines 12-19, and having both ovomucoid and conalbumin sequence elements (pINLB-MDOT-EPO vector)
- Go birds containing the EPO transgene were tested to determine expression levels of EPO in serum. Of the 62 birds tested (60 experimental birds and 2 controls), 23 were positive for EPO expression in the serum with levels > 3.0 mg/ml. The additional 27 birds had EPO levels

ranging from 0.058 ag/ml to 2.450 ng/ml. The two control birds were negative for EPO expression.

In a similar manner, serum samples were assayed by EUISA for GM-CSF production in 73  $G_0$  birds, produced as described above. Results showed that 3 of the experimental birds had serum levels of GM-CSF  $\geq$  1.0 ng/ml, while an additional 1 birds had levels ranging from 0.026 ng/ml to about 1.82 ng/ml. The remainder of the birds, and the three controls, were negative for GM-CSF production.

10. Consistent expression of β-lactamase was shown in the serum of G<sub>0</sub>, G<sub>1</sub>, and G<sub>2</sub> transgenic chickens and in the egg white of eggs from G<sub>2</sub> hens, produced according to the above-described procedures. Representative results, for example, showed serum expression in G<sub>1</sub> Hen 5657, and her G<sub>2</sub> progeny, of about 2.0 µg/ml of β-lactamase. Both G<sub>1</sub> rooster 4133 and his G<sub>2</sub> progeny demonstrated levels of β-lactamase in serum samples of about 6.0 µg/ml. β-lactamase in egg whites, from eggs of G<sub>2</sub> hens sired by rooster 4133, was expressed in levels ranging from about 0.6 µg/ml to about 1.2 µg/ml.

This declaration is made with the knowledge that willful false statements and the like are punishable by fine or imprisonment, or both (18 U.S.C. §1001) and may jeopardize the validary of the application or any patent issuing thereon. All statements made of my own knowledge are true and all statements made on information and belief are believed to be true.

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Table 1. Oligonucleotides used for IFN gene synthesis

		Prime	Γ	Prime
		1		7 11
Templa		r		r
te	Sequence of Template	1	Sequence of Primer 1	2
IFN-A	5'-	IFN-1	5'-	IFN-2
	ATGGCTTTGACCTTTGCCTTACTGGT		CCCAAGCTTTCACCATGGCTT	
	GGCTCTCCTGGT		TGACCTTTG	
	GCTGAGCTGCAAGAGCAGCTGCTCTG		CCTT-3'	
	TGGGCTGCGATCTGCCTCA-3'			
IFN-B	5'-	IFN-	5'-	IFN-
	GACCCACAGCCTGGGCAGCAGGAGG	2b	ATCTGCCTCAGACCCACAG-3'	3b
	ACCCTGATGCT			
	GCTGGCTCAGATGAGGAGAATCAGC			
	CTGTTTAGCTGCCTGAAGGATAGGCA			
	CGATTTTGGCTTT-3'			
IFN-C	5'-	IFN-	5'-	IFN-4
	CTCAAGAGGAGTTTGGCAACCAGTTT	3c	GATTTTGGCTTTCCTCAAGAG	
	CAGAAGGCTG		GAGTT-3'	
	AGACCATCCCTGTGCTGCACGAGATG			
	-3'		*	
IFN-D	5'-	IFN-	5'-	IFN-5
	TCCAGCAGATCTTTAACCTGTTTAGC	4b	GCACGAGATGATCCAGCAGA	
	ACCAAGGATAG		T-3'	
	CAGCGCTGCTTGGGATGAGACCCTGC			
	TGGATAAGTTTTACACCGAGCTGTAC			
	CAGCA-3'			

IFN-E	5'- CTGAACGATCTGGAGGCTTGCGTGAT CCAGGGCGTG GGCGTGACCGAGACCCCTCTGATGA	IFN-5	5'- ATCGTTCAGCTGCTGGTACA- 3'	IFN-6
IFN-F	AGGAGGATAGCATCCT-3' 5'- GCTGTGAGGAAGTACTTTCAGAGGAT CACCCTGTACC TGAAGGAGAAGAAGTACAGCCCTTG CGCTTGGGAAGTCGTGAGGG-3'	IFN-6	5'- CCTCACAGCCAGGATGCTAT- 3'	IFN-7
IFN-G	5'- CTGAGATCATGAGGAGCTTTAGCCTG AGCACCAACCT GCAAGAGAGAGCTTGAGGTCTAAGGAG TAA-3'		ATGATCTCAGCCCTCACGAC- 3'	IF IN-